Your Guide to Understanding Genetic Conditions

CRTAP gene

cartilage associated protein

Normal Function

The *CRTAP* gene provides instructions for making a protein called cartilage associated protein. While the specific function of this protein is not known, it plays an important role in normal bone development. Cartilage associated protein works with two other proteins, leprecan and cyclophilin B, as part of a complex that helps process certain forms of collagen. Collagens are proteins that provide strength, support, and the ability to stretch (elasticity) to many body tissues.

The complex containing cartilage associated protein modifies a protein building block (amino acid) called proline in collagen molecules. This modification, which is known as proline 3-hydroxylation, appears to be critical for the normal folding and assembly of collagen. It also may be important for releasing collagen molecules into the spaces around cells (the extracellular matrix). The secretion of collagen from cells is necessary for the proper formation of connective tissues, such as bones, tendons, and cartilage, that form the body's supportive framework.

Health Conditions Related to Genetic Changes

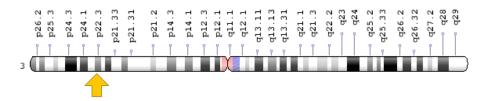
osteogenesis imperfecta

At least five mutations in the *CRTAP* gene are responsible for a rare type of osteogenesis imperfecta that is usually classified as type VII. Several of these mutations prevent cells from producing any cartilage associated protein. Without this protein, bones and other connective tissues do not form properly, leading to a very severe form of the disorder. Another mutation in the *CRTAP* gene greatly reduces the amount of cartilage associated protein produced, which disrupts the normal formation of collagen. This genetic change causes less severe signs and symptoms of osteogenesis imperfecta.

Chromosomal Location

Cytogenetic Location: 3p22.3, which is the short (p) arm of chromosome 3 at position 22.3

Molecular Location: base pairs 33,113,958 to 33,147,773 on chromosome 3 (Homo sapiens Annotation Release 108, GRCh38.p7) (NCBI)



Credit: Genome Decoration Page/NCBI

Other Names for This Gene

- cartilage-associated protein
- CASP
- CRTAP HUMAN

Additional Information & Resources

Educational Resources

- Howard Hughes Medical Institute: Genetic Mutation Explains Form of Brittle Bone Disease (October 20, 2006)
 http://www.hhmi.org/news/genetic-mutation-explains-form-brittle-bone-disease
- Molecular Biology of the Cell (fourth edition, 2002): Collagens Are the Major Proteins of the Extracellular Matrix https://www.ncbi.nlm.nih.gov/books/NBK26810/#A3551
- Molecular Cell Biology (fourth edition, 2000): Collagen: The Fibrous Proteins of the Matrix https://www.ncbi.nlm.nih.gov/books/NBK21582/
- The Cell: A Molecular Approach (second edition, 2000): Collagen fibrils (figure) https://www.ncbi.nlm.nih.gov/books/NBK9874/?rendertype=figure&id=A2050

Scientific Articles on PubMed

PubMed

https://www.ncbi.nlm.nih.gov/pubmed?term=%28%28CRTAP%5BTIAB%5D%29+OR+%28cartilage+associated+protein%5BTIAB%5D%29%29+AND+english%5Bla%5D+AND+human%5Bmh%5D+AND+%22last+3600+days%22%5Bdp%5D

OMIM

 CARTILAGE-ASSOCIATED PROTEIN http://omim.org/entry/605497

Research Resources

ClinVar

https://www.ncbi.nlm.nih.gov/clinvar?term=CRTAP%5Bgene%5D

- HGNC Gene Symbol Report http://www.genenames.org/cgi-bin/gene_symbol_report?q=data/ hgnc data.php&hgnc id=2379
- NCBI Gene https://www.ncbi.nlm.nih.gov/gene/10491
- UniProt http://www.uniprot.org/uniprot/O75718

Sources for This Summary

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Citation on PubMed: https://www.ncbi.nlm.nih.gov/pubmed/17192541

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 Citation on PubMed: https://www.ncbi.nlm.nih.gov/pubmed/12110407
- Marini JC, Cabral WA, Barnes AM, Chang W. Components of the collagen prolyl 3-hydroxylation complex are crucial for normal bone development. Cell Cycle. 2007 Jul 15;6(14):1675-81. Epub 2007 May 18. Review.
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- Morello R, Bertin TK, Chen Y, Hicks J, Tonachini L, Monticone M, Castagnola P, Rauch F, Glorieux FH, Vranka J, Bächinger HP, Pace JM, Schwarze U, Byers PH, Weis M, Fernandes RJ, Eyre DR, Yao Z, Boyce BF, Lee B. CRTAP is required for prolyl 3- hydroxylation and mutations cause recessive osteogenesis imperfecta. Cell. 2006 Oct 20;127(2):291-304.
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 Citation on PubMed: https://www.ncbi.nlm.nih.gov/pubmed/10702664
- Ward LM, Rauch F, Travers R, Chabot G, Azouz EM, Lalic L, Roughley PJ, Glorieux FH.
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Reprinted from Genetics Home Reference: https://ghr.nlm.nih.gov/gene/CRTAP

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